

#9
re-run

1600

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/769,952

DATE: 05/08/2003
 TIME: 10:08:03

Input Set : A:\US09769952.raw
 Output Set: N:\CRF4\05082003\I769952.raw

1 <110> APPLICANT: LEXICON GENETICS INCORPORATED
 2 <120> TITLE OF INVENTION: Novel Human Enzymes and Polynucleotides
 3 Encoding the Same
 4 <130> FILE REFERENCE: LEX-0118-PCT
 5 <140> CURRENT APPLICATION NUMBER: US/09/769,952
 6 <141> CURRENT FILING DATE: 2001-01-25
 7 <150> PRIOR APPLICATION NUMBER: US 60/179,000
 8 <151> PRIOR FILING DATE: 2000-01-28
 9 <160> NUMBER OF SEQ ID NOS: 15
 10 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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 13 <211> LENGTH: 831
 14 <212> TYPE: DNA
 15 <213> ORGANISM: Homo sapiens
 16 <400> SEQUENCE: 1
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 18 gtcactcgcg cttgttagctt catccggag gcagcaacgc aaggagccaa aatagttct 120
 19 ttgccgaaat gcttaatttccatatggc gcgaaatattt ttcctgaata tgccagagaaa 180
 20 attcctggtg aatccacaca gaagcttctt gaagtagcaa aggaatgcag catatatctc 240
 21 attggaggct ctatccctga agaggatgct gggaaatttat ataacacctg tgctgtgtt 300
 22 gggcctgtatg gaactttact agaaaaatgtat agaaaatgtcc atctgtttga cattgtatgtt 360
 23 cctggaaaaaa ttacatttca agaatctaaa acattgagtc cgggtgatag tttctccaca 420
 24 tttgatactc cttactgcag agtgggtctg ggcacatcgct acgacatgcg gtttgcagag 480
 25 cttgcacaaa tctacgcaca gagaggctgc cagctgttgg tatatccagg agctttat 540
 26 ctgaccactg gaccagccca ttgggagttt cttcagcgaa gccgggctgt tgataatcag 600
 27 gtgtatgtgg ccacagccctc tcctgccccgg gatgacaaag cctcctatgt tgcctgggaa 660
 28 cacagcaccc tggtgaaccc ttggggggag gttctagcca aagctggcac agaagaagca 720
 29 atcgtgtatt cagacataga cctgaagaag ctggctgaaa tacgcccagca aatccccgtt 780
 30 ttttagacaga agcgatcaga cctctatgct gtggagatga aaaagcccta a 831
 32 <210> SEQ ID NO: 2
 33 <211> LENGTH: 276
 34 <212> TYPE: PRT
 35 <213> ORGANISM: Homo sapiens
 36 <400> SEQUENCE: 2
 37 Met Thr Ser Phe Arg Leu Ala Leu Ile Gln Leu Gln Ile Ser Ser Ile
 38 1 5 10 15
 39 Lys Ser Asp Asn Val Thr Arg Ala Cys Ser Phe Ile Arg Glu Ala Ala
 40 20 25 30
 41 Thr Gln Gly Ala Lys Ile Val Ser Leu Pro Glu Cys Phe Asn Ser Pro
 42 35 40 45
 43 Tyr Gly Ala Lys Tyr Phe Pro Glu Tyr Ala Glu Lys Ile Pro Gly Glu
 44 50 55 60
 45 Ser Thr Gln Lys Leu Ser Glu Val Ala Lys Glu Cys Ser Ile Tyr Leu

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46      65          70          75          80
47 Ile Gly Gly Ser Ile Pro Glu Glu Asp Ala Gly Lys Leu Tyr Asn Thr
48           85          90          95
49 Cys Ala Val Phe Gly Pro Asp Gly Thr Leu Leu Ala Lys Tyr Arg Lys
50           100         105         110
51 Ile His Leu Phe Asp Ile Asp Val Pro Gly Lys Ile Thr Phe Gln Glu
52           115         120         125
53 Ser Lys Thr Leu Ser Pro Gly Asp Ser Phe Ser Thr Phe Asp Thr Pro
54           130         135         140
55 Tyr Cys Arg Val Gly Leu Gly Ile Cys Tyr Asp Met Arg Phe Ala Glu
56           145         150         155         160
57 Leu Ala Gln Ile Tyr Ala Gln Arg Gly Cys Gln Leu Leu Val Tyr Pro
58           165         170         175
59 Gly Ala Phe Asn Leu Thr Thr Gly Pro Ala His Trp Glu Leu Leu Gln
60           180         185         190
61 Arg Ser Arg Ala Val Asp Asn Gln Val Tyr Val Ala Thr Ala Ser Pro
62           195         200         205
63 Ala Arg Asp Asp Lys Ala Ser Tyr Val Ala Trp Gly His Ser Thr Val
64           210         215         220
65 Val Asn Pro Trp Gly Glu Val Leu Ala Lys Ala Gly Thr Glu Glu Ala
66           225         230         235         240
67 Ile Val Tyr Ser Asp Ile Asp Leu Lys Lys Leu Ala Glu Ile Arg Gln
68           245         250         255
69 Gln Ile Pro Val Phe Arg Gln Lys Arg Ser Asp Leu Tyr Ala Val Glu
70           260         265         270
71 Met Lys Lys Pro
72           275
74 <210> SEQ ID NO: 3
75 <211> LENGTH: 480
76 <212> TYPE: DNA
77 <213> ORGANISM: Homo sapiens
78 <400> SEQUENCE: 3
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80 attttctgtt tggaaacacgc ttactgcaga gtgggtctgg gcatctgcta cgacatgcgg      120
81 tttcagagc ttgcacaaaat ctacgcacag agaggctgcc agctgttggt atatccagga      180
82 gcttttaatc tgaccactgg accagcccat tgggagttac ttcaagcgaag ccgggctgtt      240
83 gataatcagg tgtatgtggc cacagcctct cctgcccggg atgacaaaagc ctcctatgtt      300
84 gcctggggac acagcacccgt ggtgaaccct tggggggagg ttctagccaa agctggcaca      360
85 gaagaagcaa tcgtgtattc agacatagac ctgaagaagc tggctgaaat acgccagcaa      420
86 atccccgttt ttagacagaaa gcgatcagac ctctatgctg tggagatgaa aaagccctaa      480
88 <210> SEQ ID NO: 4
89 <211> LENGTH: 159
90 <212> TYPE: PRT
91 <213> ORGANISM: Homo sapiens
92 <400> SEQUENCE: 4
93 Met Ser Trp Arg Ile Ser Pro Ala Thr Pro Cys Cys Arg Glu Leu Thr      15
94   1           5           10          15
95 Phe His Leu Cys Ile Phe Cys Leu Glu Thr Ala Tyr Cys Arg Val Gly      30
96           20          25

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97 Leu Gly Ile Cys Tyr Asp Met Arg Phe Ala Glu Leu Ala Gln Ile Tyr
 98 35 40 45
 99 Ala Gln Arg Gly Cys Gln Leu Leu Val Tyr Pro Gly Ala Phe Asn Leu
 100 50 55 60
 101 Thr Thr Gly Pro Ala His Trp Glu Leu Leu Gln Arg Ser Arg Ala Val
 102 65 70 75 80
 103 Asp Asn Gln Val Tyr Val Ala Thr Ala Ser Pro Ala Arg Asp Asp Lys
 104 85 90 95
 105 Ala Ser Tyr Val Ala Trp Gly His Ser Thr Val Val Asn Pro Trp Gly
 106 100 105 110
 107 Glu Val Leu Ala Lys Ala Gly Thr Glu Glu Ala Ile Val Tyr Ser Asp
 108 115 120 125
 109 Ile Asp Leu Lys Lys Leu Ala Glu Ile Arg Gln Gln Ile Pro Val Phe
 110 130 135 140
 111 Arg Gln Lys Arg Ser Asp Leu Tyr Ala Val Glu Met Lys Lys Pro
 112 145 150 155
 114 <210> SEQ ID NO: 5
 115 <211> LENGTH: 366
 116 <212> TYPE: DNA
 117 <213> ORGANISM: Homo sapiens
 118 <400> SEQUENCE: 5
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 120 ccaggagctt ttaatctgac cactggacca gcccattggg agttacttca gcgaaaggccgg 120
 121 gctgttgata atcagggtgtt tttggccaca gccttcctgt cccggatgtt caaaggctcc 180
 122 tatgttgccct ggggacacag caccgtgggtt aacccttggg gggaggttctt agccaaagct 240
 123 ggcacagaag aagcaatcgt gtattcagac atagacctga agaagctggc tgaaaatacgc 300
 124 cagcaaattcc ccgttttagt acagaagcgt tcagacctct atgctgttggatgaaaaag 360
 125 ccctaa
 127 <210> SEQ ID NO: 6
 128 <211> LENGTH: 121
 129 <212> TYPE: PRT
 130 <213> ORGANISM: Homo sapiens
 131 <400> SEQUENCE: 6
 132 Met Arg Phe Ala Glu Leu Ala Gln Ile Tyr Ala Gln Arg Gly Cys Gln
 133 1 5 10 15
 134 Leu Leu Val Tyr Pro Gly Ala Phe Asn Leu Thr Thr Gly Pro Ala His
 135 20 25 30
 136 Trp Glu Leu Leu Gln Arg Ser Arg Ala Val Asp Asn Gln Val Tyr Val
 137 35 40 45
 138 Ala Thr Ala Ser Pro Ala Arg Asp Asp Lys Ala Ser Tyr Val Ala Trp
 139 50 55 60
 140 Gly His Ser Thr Val Val Asn Pro Trp Gly Glu Val Leu Ala Lys Ala
 141 65 70 75 80
 142 Gly Thr Glu Glu Ala Ile Val Tyr Ser Asp Ile Asp Leu Lys Lys Leu
 143 85 90 95
 144 Ala Glu Ile Arg Gln Gln Ile Pro Val Phe Arg Gln Lys Arg Ser Asp
 145 100 105 110
 146 Leu Tyr Ala Val Glu Met Lys Lys Pro
 147 115 120

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149 <210> SEQ ID NO: 7
 150 <211> LENGTH: 507
 151 <212> TYPE: DNA
 152 <213> ORGANISM: Homo sapiens
 153 <400> SEQUENCE: 7

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155	attttctgtt	tggaaacagc	ttactgcaga	gtgggtctgg	gcatctgcta	cgacatgcgg	120
156	tttcagagc	ttgcacaaat	ctacgcacag	agaggctgcc	agctgttgt	atateccagga	180
157	gttttaatc	tgaccactgg	accagccat	tggagttac	ttcagcgaag	ccgggctgtt	240
158	gataatcagg	tgtatgtggc	cacagcctct	cctgcccggg	atgacaaaagc	ctcctatgtt	300
159	gcctggggac	acagcaccgt	ggtgaaccct	tggggggagg	ttctagccaa	agctggcaca	360
160	gaagaagcaa	tcgtgtattc	agacatagac	ctgaagaagc	tggctgaaat	acgcccagcaa	420
161	atccccgtt	ttagacagaa	gcgaaatatt	ttcctgaata	tgcagagaaa	attcctggtg	480
162	aatccacaca	gaagcttct	gaagtag				507

164 <210> SEQ ID NO: 8
 165 <211> LENGTH: 168
 166 <212> TYPE: PRT
 167 <213> ORGANISM: Homo sapiens
 168 <400> SEQUENCE: 8

169	Met	Ser	Trp	Arg	Ile	Ser	Pro	Ala	Thr	Pro	Cys	Cys	Arg	Glu	Leu	Thr
170	1				5				10					15		
171	Phe	His	Leu	Cys	Ile	Phe	Cys	Leu	Glu	Thr	Ala	Tyr	Cys	Arg	Val	Gly
172									20			25		30		
173	Leu	Gly	Ile	Cys	Tyr	Asp	Met	Arg	Phe	Ala	Glu	Leu	Ala	Gln	Ile	Tyr
174							35		40			45				
175	Ala	Gln	Arg	Gly	Cys	Gln	Leu	Leu	Val	Tyr	Pro	Gly	Ala	Phe	Asn	Leu
176							50		55			60				
177	Thr	Thr	Gly	Pro	Ala	His	Trp	Glu	Leu	Leu	Gln	Arg	Ser	Arg	Ala	Val
178							65		70		75		80			
179	Asp	Asn	Gln	Val	Tyr	Val	Ala	Thr	Ala	Ser	Pro	Ala	Arg	Asp	Asp	Lys
180							85		90			95				
181	Ala	Ser	Tyr	Val	Ala	Trp	Gly	His	Ser	Thr	Val	Val	Asn	Pro	Trp	Gly
182							100		105			110				
183	Glu	Val	Leu	Ala	Lys	Ala	Gly	Thr	Glu	Glu	Ala	Ile	Val	Tyr	Ser	Asp
184							115		120			125				
185	Ile	Asp	Leu	Lys	Lys	Leu	Ala	Glu	Ile	Arg	Gln	Gln	Ile	Pro	Val	Phe
186							130		135			140				
187	Arg	Gln	Lys	Arg	Asn	Ile	Phe	Leu	Asn	Met	Gln	Arg	Lys	Phe	Leu	Val
188							145		150		155		160			
189	Asn	Pro	His	Arg	Ser	Phe	Leu	Lys								
190						165										

192 <210> SEQ ID NO: 9
 193 <211> LENGTH: 393
 194 <212> TYPE: DNA
 195 <213> ORGANISM: Homo sapiens
 196 <400> SEQUENCE: 9

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198	ccaggagctt	ttaatctgac	cactggacca	gcccatgggg	agttacttca	gcgaagccgg	120
199	gctgttata	atcaggtgta	tgtggccaca	gcctctcctg	cccggtatga	caaagcctcc	180

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200 tatgttgcct gggcacacag caccgtggc aacccttggg gggagggtct agccaaagct 240
 201 ggcacagaag aagcaatcg gtattcagac atagacctga agaagctggc taaaatacgc 300
 202 cagcaaattcc cggttttag acagaagcga aatattttcc tgaatatgca gagaaaattc 360
 203 ctggtaatc cacacagaag cttctgaa tag
 205 <210> SEQ ID NO: 10
 206 <211> LENGTH: 130
 207 <212> TYPE: PRT
 208 <213> ORGANISM: Homo sapiens
 209 <400> SEQUENCE: 10
 210 Met Arg Phe Ala Glu Leu Ala Gln Ile Tyr Ala Gln Arg Gly Cys Gln 210
 211 1 5 10 15
 212 Leu Leu Val Tyr Pro Gly Ala Phe Asn Leu Thr Thr Gly Pro Ala His 210
 213 20 25 30
 214 Trp Glu Leu Leu Gln Arg Ser Arg Ala Val Asp Asn Gln Val Tyr Val 210
 215 35 40 45
 216 Ala Thr Ala Ser Pro Ala Arg Asp Asp Lys Ala Ser Tyr Val Ala Trp 210
 217 50 55 60
 218 Gly His Ser Thr Val Val Asn Pro Trp Gly Glu Val Leu Ala Lys Ala 210
 219 65 70 75 80
 220 Gly Thr Glu Glu Ala Ile Val Tyr Ser Asp Ile Asp Leu Lys Lys Leu 210
 221 85 90 95
 222 Ala Glu Ile Arg Gln Gln Ile Pro Val Phe Arg Gln Lys Arg Asn Ile 210
 223 100 105 110
 224 Phe Leu Asn Met Gln Arg Lys Phe Leu Val Asn Pro His Arg Ser Phe 210
 225 115 120 125
 226 Leu Lys
 227 130
 229 <210> SEQ ID NO: 11
 230 <211> LENGTH: 459
 231 <212> TYPE: DNA
 232 <213> ORGANISM: Homo sapiens
 233 <400> SEQUENCE: 11
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 235 gtcactcgcg cttgtagctt catccgggag gcagcaacgc aaggagccaa aatagtttct 120
 236 ttgcggaat gcttaattc tccatatggc gcgaaatatt ttcctgaata tgcagagaaa 180
 237 attcctggtg aatccacaca gaagtttct gaagtagccaa aggaatgcag catatatctc 240
 238 attggaggct ctatccctga agaggatgtt gggaaattat ataacacctg tgctgtgttt 300
 239 gggctgtatg gaactttact agcaaagtat agaaagatcc atctgtttga cattgatgtt 360
 240 cctggaaaaa ttacatttca agaatctaaa acattgagtc cgggtgatag tttctccaca 420
 241 tttgatactc gtatgtacca gataagtttgc cctcttttag 459
 243 <210> SEQ ID NO: 12
 244 <211> LENGTH: 152
 245 <212> TYPE: PRT
 246 <213> ORGANISM: Homo sapiens
 247 <400> SEQUENCE: 12
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 249 1 5 10 15
 250 Lys Ser Asp Asn Val Thr Arg Ala Cys Ser Phe Ile Arg Glu Ala Ala 248
 251 20 25 30

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/769,952

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Input Set : A:\US09769952.raw

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